

GenCore version 4.5  
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OM nucleic - nucleic search, using sw model

Run on: November 25, 2000, 04:35:30 ; Search time 84.1 Seconds  
(without alignments)  
847.024 Million cell updates/sec

Title: US-09-373-230-1  
Perfect score: 471  
Sequence: 1 AACCTTGCCGACTTCACTG.....TCACTTAACCTTACATCAAGT 471

Scoring table: IDENTITY\_NUC  
Gapop 10.0 , Gapext 1.0

Searched: 262060 seqs, 75620727 residues

Total number of hits satisfying chosen parameters: 524120

Minimum DB seq length: 0  
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%  
Maximum Match 100%  
Listing first 45 summaries

Database : Issued Patents.NA.\*  
1: /cgn2\_6/ptodata/1/ina/5A.COMB.seq.\*  
2: /cgn2\_6/ptodata/1/ina/5B.COMB.seq.\*  
3: /cgn2\_6/ptodata/1/ina/5C.COMB.seq.\*  
4: /cgn2\_6/ptodata/1/ina/5D.COMB.seq.\*  
5: /cgn2\_6/ptodata/1/ina/6.COMB.seq.\*  
6: /cgn2\_6/ptodata/1/ina/PCRUS.COMB.seq.\*  
7: /cgn2\_6/ptodata/1/ina/backfiles1.seq.\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	ID	Description
1	470.6	99.9	471	4	US-08-502-535B-1
2	470.6	99.9	471	4	US-08-908-005A-1
3	247.4	52.5	579	3	US-08-896-605A-7
4	247.4	52.5	579	3	US-08-896-501A-5
5	247.4	52.5	1120	5	US-08-884-324-2
6	127.8	27.1	11464	5	US-08-884-324-13
7	127.8	27.1	28994	5	US-08-884-324-14
8	122	25.9	2167	5	US-08-884-324-7
9	75.6	16.1	134	5	US-08-884-324-4
10	51.4	10.9	135	5	US-08-884-324-3
11	38.4	9.2	7218	1	US-08-232-463-14
12	36.8	7.8	4731	5	US-08-488-706-2
13	36.8	7.8	4731	5	US-08-772-270A-9
14	34.4	7.3	10614	1	US-08-135-511-35
15	34.4	7.3	10614	2	US-08-187-453-35
16	33.8	7.2	8920	3	US-08-446-855A-1
17	33.2	7.0	4467	2	US-08-565-907A-1
18	33.2	7.0	4467	4	US-08-910-551B-1
19	33.2	7.0	4467	4	US-08-909-425A-1
20	32.6	6.9	4821	1	US-08-053-614-3
21	32.6	6.9	4821	2	US-08-316-397B-3
22	32.6	6.9	4821	3	US-09-034-306-3
23	32.6	6.9	4821	6	PCT-US93-09782-3
24	32.4	6.9	1326	5	US-09-100-391-1
25	32.4	6.9	3095	7	5231168-1
26	32	6.8	1431	2	US-08-451-715A-11
27					Sequence 1, Appli
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33	29.8	6.3	5035	2	US-08-618-392C-3	Sequence 1, Appli
34	29.6	6.3	26700	2	US-08-472-217-1	Sequence 1, Appli
35	29.6	6.3	26700	5	US-08-488-159-5	Sequence 1, Appli
36	29.6	6.3	26700	5	US-08-760-534A-1	Sequence 1, Appli
37	29.2	6.2	1600	2	US-09-028-819-14	Sequence 1, Appli
38	29.2	6.2	1600	5	US-08-038-768A-1	Sequence 1, Appli
39	29	6.2	5035	1	US-08-391-743A-1	Sequence 1, Appli
40	28.8	6.1	1837	4	US-08-909-965C-10	Sequence 1, Appli
41	28.8	6.1	2295	1	US-08-375-300-3	Sequence 1, Appli
42	28.8	6.1	2295	5	US-09-177-431-3	Sequence 3, Appli
43	28.8	6.1	2295	6	PCT-US95-16930-3	Sequence 3, Appli
44	28.8	6.1	3634	5	US-09-166-186-1	Sequence 1, Appli
45	28.8	6.1	3792	4	US-08-992-334-1	Sequence 1, Appli

ALIGNMENTS

RESULT 1  
US-08-502-535B-1  
; Sequence 1, Application US/08502535B  
; Patent No. 5912324  
; GENERAL INFORMATION:  
; APPLICANT: OKAMURA, Haruki  
; APPLICANT: TANIMOTO, Tadao  
; APPLICANT: TORIGOE, Kakuji  
; APPLICANT: KONIKATA, Toshio  
; APPLICANT: TANIGUCHI, Mutsuko  
; APPLICANT: KOHNO, Keizo  
; APPLICANT: KURIMOTO, Masashi  
; TITLE OF INVENTION: IFN-BETA PRODUCTION INDUCING PROTEIN AND  
; TITLE OF INVENTION: MONOCLONAL ANTIBODY OF THE SAME  
; NUMBER OF SEQUENCES: 9  
; CORRESPONDENCE ADDRESS:  
; ADDRESSEE: BROWDY AND NEIMARK  
; STREET: 419 Seventh Street, N.W., Suite 300  
; CITY: Washington  
; STATE: D.C.  
; COUNTRY: USA  
; ZIP: 20004  
; COMPUTER READABLE FORM:  
; MEDIUM TYPE: Floppy disk  
; COMPUTER: IBM PC compatible  
; OPERATING SYSTEM: PC-DOS/MS-DOS  
; SOFTWARE: PatentIn Release #1.0, Version #1.30  
; CURRENT APPLICATION DATA:  
; APPLICATION NUMBER: US/08/502.535B  
; FILING DATE: 14-JUL-1995  
; CLASSIFICATION: 530  
; PRIOR APPLICATION DATA:  
; APPLICATION NUMBER: JP 184162/1994  
; FILING DATE: 14-JUL-1994  
; PRIOR APPLICATION NUMBER:  
; APPLICATION DATA:  
; FILING DATE: 10-FEB-1995  
; ATTORNEY/AGENT INFORMATION:  
; NAME: BROWDY, Roger L.  
; REGISTRATION NUMBER: 25,618  
; REFERENCE/DOCKET NUMBER: OKAMURA-2  
; TELECOMMUNICATION INFORMATION:  
; TELEPHONE: 202-628-5197  
; TELEFAX: 202-737-3528  
; INFORMATION FOR SEQ ID NO: 1:  
; SEQUENCE CHARACTERISTICS:  
; LENGTH: 471 base pairs  
; TYPE: nucleic acid  
; STRANDEDNESS: single

;; TOPOLOGY: linear  
;; MOLECULE TYPE: cDNA  
;; FEATURE:

;; NAME/KEY: CDS

;; LOCATION: 1..471

;; OTHER INFORMATION: /note= xaa in position 70 is Met or Thr

US-08-502-535B-1

Query Match 99.9%; Score 470.6; DB 4; Length 471;  
Best Local Similarity 100.0%; Pred. No. 2.4e-132;  
Matches 471; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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Db 1 AACTTGGCCGACTTCACTGTACAAACCGCAGTATACCGGAATATAATGACCAGTTCTC 60  
Qy 61 TTCGTTGACAAAACACAGCCTGTCTTCGAGGATATGACTGATTTGATCAAAAGTGCCAGT 120  
Db 61 TTCGTTGACAAAACACAGCCTGTCTTCGAGGATATGACTGATTTGATCAAAAGTGCCAGT 120  
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Qy 181 GTGACCTCTCTGTGAAGGATAGTAAAYGTCTACCTCTCTCTGTAGAAACAAGATCATT 240  
Db 181 GTGACCTCTCTGTGAAGGATAGTAAAYGTCTACCTCTCTCTGTAGAAACAAGATCATT 240  
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Db 301 TTTGAGAACTGTTCAGGACACAAAGATGGAGTTTGAATCTTCACTGTATGAAGGA 360  
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Db 361 CACTTCTCTGCTCCAAAAGGAGATGCTTTTCAAACTCATTCGAAAAAAGGAT 420  
Qy 421 GAAATGGGGATAAATCTGTAATCTTCACTCTCACTCACTTACATCAAAAGT 471  
Db 421 GAAATGGGGATAAATCTGTAATCTTCACTCTCACTCACTTACATCAAAAGT 471

## RESULT 2

US-08-908-005A-1  
; Sequence 1, Application US/08908005A

; Patent No. 5914253

; GENERAL INFORMATION:

; APPLICANT: OKAMURA, Haruki

; APPLICANT: TANIMOTO, Tadao

; APPLICANT: TORIGOE, Kakuji

; APPLICANT: KUNIKATA, Toshio

; APPLICANT: TANIGUCHI, Mutsuko

; APPLICANT: KOHNO, Keizo

; APPLICANT: KURIMOTO, Masashi

; TITLE OF INVENTION: IFN-BETA PRODUCTION INDUCING PROTEIN AND

; TITLE OF INVENTION: MONOCLONAL ANTIBODY OF THE SAME

; NUMBER OF SEQUENCES: 9

; CORRESPONDENCE ADDRESS:

; ADDRESSEE: BROWDY, Roger L.

; STREET: 419 Seventh Street, N.W., Suite 300

; CITY: Washington

; STATE: D.C.

; COUNTRY: USA

; ZIP: 20004

; COMPUTER READABLE FORM:

; MEDIUM TYPE: Floppy disk

; COMPUTER: IBM PC compatible

; OPERATING SYSTEM: PC-DOS/MS-DOS

; SOFTWARE: Patentin Release #1.0, Version #1.30

;; CURRENT APPLICATION DATA:  
;; APPLICATION NUMBER: US/08/908.005A  
;; FILING DATE: 11-AUG-1997  
;; CLASSIFICATION: 530  
;; PRIOR APPLICATION DATA:  
;; APPLICATION NUMBER: US 08/502,535  
;; FILING DATE: 14-JUL-1995  
;; PRIOR APPLICATION DATA:  
;; APPLICATION NUMBER: JP 184162/1994  
;; FILING DATE: 14-JUL-1994  
;; PRIOR APPLICATION DATA:  
;; APPLICATION NUMBER: JP 45057/1995  
;; FILING DATE: 10-FEB-1995  
;; ATTORNEY/AGENT INFORMATION:  
;; NAME: BROWDY, Roger L.  
;; REGISTRATION NUMBER: 25,618  
;; REFERENCE/DOCKET NUMBER: OKAMURA-2A  
;; TELECOMMUNICATION INFORMATION:  
;; TELEPHONE: 202-628-5197  
;; TELEFAX: 202-737-3528  
;; INFORMATION FOR SEQ ID NO: 1:  
;; SEQUENCE CHARACTERISTICS:  
;; LENGTH: 471 base pairs  
;; TYPE: nucleic acid  
;; STRANDEDNESS: single  
;; TOPOLOGY: linear  
;; MOLECULE TYPE: cDNA  
;; FEATURE:  
;; NAME/KEY: CDS  
;; LOCATION: 1..471  
;; OTHER INFORMATION: /note= xaa in position 70 is Met or Thr  
US-08-908-005A-1

Query Match 99.9%; Score 470.6; DB 4; Length 471;  
Best Local Similarity 100.0%; Pred. No. 2.4e-132;  
Matches 471; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 AACTTGGCCGACTTCACTGTACAAACCGCAGTAAATACGGAATATAATGACCAGTTCTC 60  
Db 1 AACTTGGCCGACTTCACTGTACAAACCGCAGTAAATACGGAATATAATGACCAGTTCTC 60  
Qy 61 TTCGTTGACAAAACACAGCCTGTGTTCGAGGATATGACTGATTTGATCAAAAGTGCCAGT 120  
Db 61 TTCGTTGACAAAACACAGCCTGTGTTCGAGGATATGACTGATTTGATCAAAAGTGCCAGT 120  
Qy 121 GAACCCGACAGCAGCATATAATATACATGTACAAAGACAGTGAAGTAAAGGAGTGGCT 180  
Db 121 GAACCCGACAGCAGCATATAATATACATGTACAAAGACAGTGAAGTAAAGGAGTGGCT 180  
Qy 181 GTGACCTCTCTGTGAAGGATAGTAAAYGTCTACCTCTCTCTGTAAAGAACAGATCAATT 240  
Db 181 GTGACCTCTCTGTGAAGGATAGTAAAYGTCTACCTCTCTCTGTAAAGAACAGATCAATT 240  
Qy 241 TCCTTTGAGGAATGGATCCACCTGAAATATTTGATGATATACAAAGTGTATCATATTC 300  
Db 241 TCCTTTGAGGAATGGATCCACCTGAAATATTTGATGATATACAAAGTGTATCATATTC 300  
Qy 301 TTTGAGAACTGTTCAGGACACAAAGATGGAGTTTGAATCTTCACTGTATGAAGGA 360  
Db 301 TTTGAGAACTGTTCAGGACACAAAGATGGAGTTTGAATCTTCACTGTATGAAGGA 360  
Qy 361 CACTTCTCTGCTCCAAAAGGAGATGCTTTTCAAACTCATTCGAAAAAAGGAT 420  
Db 361 CACTTCTCTGCTCCAAAAGGAGATGCTTTTCAAACTCATTCGAAAAAAGGAT 420  
Qy 421 GAAATGGGGATAAATCTGTAATCTTCACTCTCACTCACTTACATCAAAAGT 471  
Db 421 GAAATGGGGATAAATCTGTAATCTTCACTCTCACTCACTTACATCAAAAGT 471

## RESULT 3

US-08-896-605A-7

; Sequence 7, Application US/08896605A  
; Patent No. 5879942  
; GENERAL INFORMATION:  
; APPLICANT: TANIMOTO, Tadao  
; APPLICANT: KURIMOTO, Masashi  
; TITLE OF INVENTION: PROCESSING ENZYME FOR POLYPEPTIDE  
; NUMBER OF SEQUENCES: 9  
; CORRESPONDENCE ADDRESS:  
; ADDRESSEE: BROWDY AND NEIMARK  
; STREET: 419 Seventh Street, N.W., Suite 300  
; CITY: Washington  
; STATE: D.C.  
; COUNTRY: USA  
; ZIP: 20004  
; COMPUTER READABLE FORM:  
; MEDIUM TYPE: Floppy disk  
; COMPUTER: IBM PC compatible  
; OPERATING SYSTEM: PC-DOS/MS-DOS  
; SOFTWARE: Patent In Release #1.0, Version #1.30  
; CURRENT APPLICATION DATA:  
; APPLICATION NUMBER: US/08/896,605A  
; FILING DATE: 18 JULY 1997  
; PRIOR APPLICATION DATA:  
; APPLICATION NUMBER: JP 207,691/1996  
; FILING DATE: 19-JUL-1996  
; APPLICATION DATA:  
; APPLICATION NUMBER: JP 156,062/1997  
; FILING DATE: 30-MAY-1997  
; ATTORNEY/AGENT INFORMATION:  
; NAME: BROWDY, Roger L.  
; REGISTRATION NUMBER: 25,618  
; REFERENCE/DOCKET NUMBER: TANIMOTO=2  
; TELECOMMUNICATION INFORMATION:  
; TELEPHONE: 202-628-5197  
; TELEFAX: 202-737-3528  
; INFORMATION FOR SEQ ID NO: 7:  
; SEQUENCE CHARACTERISTICS:  
; LENGTH: 579 base pairs  
; TYPE: nucleic acid  
; STRANDEDNESS: single  
; TOPOLOGY: linear  
; MOLECULE TYPE: cdna to mRNA  
; FEATURE:  
; NAME/KEY: leader peptide  
; LOCATION: 1..108  
; IDENTIFICATION METHOD: S  
; NAME/KEY: mat peptide  
; LOCATION: 109..579  
; IDENTIFICATION METHOD: S  
; US-08-896-605A-7

Query Match 52.5%; Score 247.4; DB 3; Length 579;  
Best Local Similarity 73.1%; Pred. No. 2.4e-65;  
Matches 343; Conservative 2; Mismatches 118; Indels 6; Gaps 2;  
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QY 62 TCGTTTGCAAAAGACA---GGCTGTGTTGAGGATATGACTGATATTGATCAAAAGTGCCA 118  
Db 170 TCATTGACCAAGGAATCGGCCTCTATTTCAGATATGACTGCTGCTGAGAGATA 229  
QY 119 GTGAGCCCGAGCAGCTGATATATACATGACAAAGCAGTGAAGTAAAGGACTGG 178  
Db 230 ATGCACCCCGGACCATTTATTAATAGTAATGATAGATAGCCACGCTAGAGGTGG 289  
QY 179 CTGTGACCCCTCTGTGGAAGGATAGTAAAYGTCTACCCCTCTCTGTGAAGCAAGATCA 238  
Db 290 CTGTAACTATCTCTGTGAAGTGTGAGAAATTTCAAYTCTCTCTGTGAGAAACAAATTA 349  
QY 239 TTTCCTTTGAGGAATGGATCCAGCTCGAANAATATTGATGATATACAAAGTGATCTCATAT 298

Db 350 TTTCCTTTAAGGAATGAATCTCTCTGATAACATCAAGGATACAAAAAGTGACATCATAT 409  
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Db 410 TCTTTTCAGAGAAGTGTCCCGAGCAGCATGATAATAGATGCAATTTGAATCTTTCATCATACG 469  
QY 356 AAGGACACTTCTTCTGCTTCCAAAGGAAGATGATGCTTTTCAAACTCATTTCTGAAAAAAA 415  
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RESULT 4  
US-08-896-501A-5  
; Sequence 5, Application US/08896501A  
; Patent No. 5891663  
; GENERAL INFORMATION:  
; APPLICANT: TANIMOTO, Tadao  
; APPLICANT: KURIMOTO, Masashi  
; TITLE OF INVENTION: PROCESS FOR PRODUCING POLYPEPTIDE  
; NUMBER OF SEQUENCES: 9  
; CORRESPONDENCE ADDRESS:  
; ADDRESSEE: BROWDY AND NEIMARK  
; STREET: 419 Seventh Street, N.W., Suite 300  
; CITY: Washington  
; STATE: D.C.  
; COUNTRY: USA  
; ZIP: 20004  
; COMPUTER READABLE FORM:  
; MEDIUM TYPE: Floppy disk  
; COMPUTER: IBM PC compatible  
; OPERATING SYSTEM: PC-DOS/MS-DOS  
; SOFTWARE: Patent In Release #1.0, Version #1.30  
; CURRENT APPLICATION DATA:  
; APPLICATION NUMBER: US/08/896,501A  
; FILING DATE: 18-JUL-1997  
; PRIOR APPLICATION DATA:  
; APPLICATION NUMBER: JP 213,267/1996  
; FILING DATE: 25-JUL-1996  
; APPLICATION DATA:  
; APPLICATION NUMBER: JP 31,474/1997  
; FILING DATE: 31-JAN-1997  
; ATTORNEY/AGENT INFORMATION:  
; NAME: BROWDY, Roger L.  
; REGISTRATION NUMBER: 25,618  
; REFERENCE/DOCKET NUMBER: TANIMOTO=3  
; TELECOMMUNICATION INFORMATION:  
; TELEPHONE: 202-628-5197  
; TELEFAX: 202-737-3528  
; INFORMATION FOR SEQ ID NO: 5:  
; SEQUENCE CHARACTERISTICS:  
; LENGTH: 579 base pairs  
; TYPE: nucleic acid  
; STRANDEDNESS: single  
; TOPOLOGY: linear  
; MOLECULE TYPE: cdna to mRNA  
; FEATURE:  
; NAME/KEY: leader peptide  
; LOCATION: 1..108  
; IDENTIFICATION METHOD: S  
; NAME/KEY: mat peptide  
; LOCATION: 109..579  
; IDENTIFICATION METHOD: S  
; US-08-896-501A-5

Query Match 52.5%; Score 247.4; DB 3; Length 579;  
Best Local Similarity 73.1%; Pred. No. 2.4e-65;  
Matches 343; Conservative 2; Mismatches 118; Indels 6; Gaps 2;



NUMBER OF SEQUENCES: 35  
CORRESPONDENCE ADDRESS:  
ADDRESSEE: BROWDY AND NEIMARK  
STREET: 419 Seventh Street, N.W., Suite 300  
CITY: Washington  
STATE: D.C.  
COUNTRY: USA  
ZIP: 20004  
COMPUTER READABLE FORM:  
MEDIUM TYPE: Floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: Patent In Release #1.0, Version #1.30  
CURRENT APPLICATION DATA:  
APPLICANT: BROWDY AND NEIMARK  
ATTORNEY/AGENT INFORMATION:  
NAME: BROWDY, Roger L.  
REGISTRATION NUMBER: 25,618  
REFERENCE/DOCKET NUMBER: OKURA=1  
TELEPHONE: 202-628-5197  
TELEFAX: 202-737-3528  
INFORMATION FOR SEQ ID NO: 13:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 11464 base pairs  
TYPE: nucleic acid  
STRANDEDNESS: double  
TOPOLOGY: linear  
MOLECULE TYPE: Genomic DNA  
ORIGINAL SOURCE:  
ORGANISM: human  
TISSUE TYPE: placenta  
FEATURE:  
NAME/KEY: 5'UTR  
LOCATION: 1..3  
IDENTIFICATION METHOD: E  
NAME/KEY: leader peptide  
LOCATION: 4..82  
IDENTIFICATION METHOD: S  
NAME/KEY: intron  
LOCATION: 83..1453  
IDENTIFICATION METHOD: E  
NAME/KEY: leader peptide  
LOCATION: 1454..1465  
IDENTIFICATION METHOD: S  
NAME/KEY: intron  
LOCATION: 1466..14848  
IDENTIFICATION METHOD: E  
NAME/KEY: leader peptide  
LOCATION: 4849..4865  
IDENTIFICATION METHOD: S  
NAME/KEY: mat peptide  
LOCATION: 4866..4983  
IDENTIFICATION METHOD: S  
NAME/KEY: intron  
LOCATION: 4984..6317  
IDENTIFICATION METHOD: E  
NAME/KEY: mat peptide  
LOCATION: 6318..6451  
IDENTIFICATION METHOD: S  
NAME/KEY: intron  
LOCATION: 6452..11224  
IDENTIFICATION METHOD: E  
NAME/KEY: mat peptide  
LOCATION: 11225..11443  
IDENTIFICATION METHOD: S  
NAME/KEY: 3'UTR  
LOCATION: 11444..11464

IDENTIFICATION METHOD: E  
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Query Match 27.1% Score 127.8; DB 5; Length 11464;  
Best Local Similarity 73.3%; Pred. No. 6.7e-29;  
Matches 178; Conservative 0; Mismatches 62; Indels 3; Gaps 1;  
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DB 11320 ATCTTCATCATACGAGGATACCTTTCTAGCTTGTGAAAAGAGAGAGACCTTTTAACT 11379  
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QY 462 ACA 464  
DB 11440 AGA 11442  
RESULT 7  
US-08-884-324-14  
; Sequence 14, Application US/08884324  
; Patent No. 6060283  
; GENERAL INFORMATION:  
; APPLICANT: Takanori OKURA  
; APPLICANT: Kakuji TORIGOE  
; APPLICANT: Masahiro KURIMOTO  
; TITLE OF INVENTION: GENOMIC DNA ENCODING A POLYPEPTIDE CAPABLE  
; OF INDUCING THE PRODUCTION OF INTERFERON-  
; NUMBER OF SEQUENCES: 35  
; CORRESPONDENCE ADDRESS:  
; ADDRESSEE: BROWDY AND NEIMARK  
; STREET: 419 Seventh Street, N.W., Suite 300  
; CITY: Washington  
; STATE: D.C.  
; COUNTRY: USA  
; ZIP: 20004  
; COMPUTER READABLE FORM:  
; MEDIUM TYPE: Floppy disk  
; OPERATING SYSTEM: PC-DOS/MS-DOS  
; SOFTWARE: Patent In Release #1.0, Version #1.30  
; CURRENT APPLICATION DATA:  
; APPLICANT: BROWDY AND NEIMARK  
; FILING DATE: 27-JUN-1996  
; PRIOR APPLICATION DATA:  
; APPLICATION NUMBER: JP 185,305/96  
; FILING DATE: 27-JUN-1996  
; ATTORNEY/AGENT INFORMATION:  
; NAME: BROWDY, Roger L.  
; REGISTRATION NUMBER: 25,618  
; REFERENCE/DOCKET NUMBER: OKURA=1  
; TELEPHONE: 202-628-5197  
; TELEFAX: 202-737-3528  
; INFORMATION FOR SEQ ID NO: 14:  
; SEQUENCE CHARACTERISTICS:  
; LENGTH: 28994 base pairs  
; TYPE: nucleic acid  
; STRANDEDNESS: double  
; TOPOLOGY: linear  
; MOLECULE TYPE: Genomic DNA









GENERAL INFORMATION:  
APPLICANT: MacInnes, Janet  
APPLICANT: Ricciatti, Paul  
APPLICANT: Mallard, Bonnie  
APPLICANT: Rosendal, Soren  
TITLE OF INVENTION: NOVEL BACTERIAL PREPARATIONS, METHOD FOR  
PRODUCING SAME, AND THEIR USE AS VACCINES  
NUMBER OF SEQUENCES: 14  
CORRESPONDENCE ADDRESS:  
ADDRESSEE: Bereskin & Parr  
STREET: 40 King Street West  
CITY: Toronto  
STATE: Ontario  
COUNTRY: Canada  
ZIP: M5H 3Y2

COMPUTER READABLE FORM:  
MEDIUM TYPE: Floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: PatentIn Release #1.0, Version #1.30  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/08/772,270A  
FILING DATE: December 23, 1996  
CLASSIFICATION: 424  
ATTORNEY/AGENT INFORMATION:  
NAME: Gravelle, Michelle  
REGISTRATION NUMBER: 40,261  
REFERENCE/DOCKET NUMBER: 6580-81  
TELEPHONE: (416) 364-7311  
TELEFAX: (416) 361-1398  
INFORMATION FOR SEQ ID NO: 9:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 4731 base pairs  
TYPE: nucleic acid  
STRANDEDNESS: single  
TOPOLOGY: linear  
MOLECULE TYPE: DNA (genomic)  
ORIGINAL SOURCE:  
ORGANISM: Actinobacillus pleuropneumoniae

US-08-772-270A-9  
Query Match 7.8%; Score 36.8; DB 5; Length 4731;  
Best Local Similarity 47.4%; Pred. No. 0.091;  
Matches 110; Conservative 0; Mismatches 122; Indels 0; Gaps 0;

QY 211 TCTACCTCTCTGTGTAAGCAAGATCATTTCTTGTGAGGAATGGATCCACCTGAARAT 270  
DB 591 TCTATCTCTAGCAAAAAGTACTGAAAATCATATATTTAAAGGAGGAAAATAACAA 650  
QY 271 ATTGATGATATCAAGATGATCTATATTTTCAGAAACGTGTTCCAGGACACACACAG 330  
DB 651 AAAAGTAGCTGAAAAGATTTCTCAGTATGAGCAAGATTTATACAGCTCTACAAT 710  
QY 331 ATGAGATTTGAATCTTCTACTGTATGAGGACACTTTCTGTTGCTGCAAAAAGGAGATGAT 390  
DB 711 AATATCTTTAAATGATCAATATATATAAGGAGACTCTTTTATGTCAAAATACACTTTGTC 770  
QY 391 GCTTTCAAACTATTCTGAAAAGGAGATGAAAATGGGATAAATCTGTAA 442  
DB 771 ATCATTAATAATCTGCTTCAACAAGGATTTGAAAATGGGAAAACAAAGTTA 822

RESULT 14  
US-08-135-511-35/c  
Sequence 35, Application US/08135511  
Patent No. 5558999  
GENERAL INFORMATION:  
APPLICANT: Chiang, John  
TITLE OF INVENTION: Cholesterol 7a-Hydroxylase Gene  
TITLE OF INVENTION: Regulatory Elements and Methods for Using Them  
NUMBER OF SEQUENCES: 35

CORRESPONDENCE ADDRESS:  
ADDRESSEE: Foley & Lardner  
STREET: 3000 K Street, N.W., Suite 500  
CITY: Washington, D.C.  
COUNTRY: USA  
ZIP: 20007-5109  
COMPUTER READABLE FORM:  
MEDIUM TYPE: Floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: PatentIn Release #1.0, Version #1.25  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/08/135,511  
FILING DATE: 13-OCT-1993  
CLASSIFICATION: 435  
ATTORNEY/AGENT INFORMATION:  
NAME: SANDERCOCK, Colin G.  
REGISTRATION NUMBER: 31,298  
REFERENCE/DOCKET NUMBER: 18748/175  
TELEPHONE: (202)672-5300  
TELEFAX: (202)672-5399  
TELEX: 904136  
INFORMATION FOR SEQ ID NO: 35:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 10614 base pairs  
TYPE: nucleic acid  
STRANDEDNESS: single  
TOPOLOGY: linear  
MOLECULE TYPE: DNA (genomic)  
US-08-135-511-35

Query Match 7.3%; Score 34.4; DB 1; Length 10614;  
Best Local Similarity 46.6%; Pred. No. 0.68; Indels 0; Gaps 0;  
Matches 110; Conservative 0; Mismatches 126; Indels 0; Gaps 0;

QY 229 AACAAAGATCATTTCTTTGAGAAATGGATCCACCTGAAAATATTGATGATATACAAAGT 288  
DB 10250 AACCTTACGAAGACCTTTGAGTCATGATTTTGAATATTTTGTGAGTTCTTGAACA 10191  
QY 289 GATCTCATATTTCTTCAGAAACGTGTTCCAGGACACACAAAGATGGAGTTTGAATCTTCA 348  
DB 10190 AATCAGATAAATATTTTAACTATCATGATGATGAATAAACAATCTATTTCAAAGGTTA 10131  
QY 349 CTGTATGAGGACACTTTCTTGTGTTCCAAAAGGAAGATGATGCTTTCAAACATCTCTG 408  
DB 10130 AAGCACAATGTTCTTATTTTCTGTTGAAAGGTAGTAAATTTTAAAGTTACTGGTTT 10071  
QY 409 AAAAAAGGATGAAATGGGATAAATCTGTAATGTTCACTCTCACTAACTTACA 464  
DB 10070 AATATTTACTTATATTTCTAGTATAAGATAAAGATAAATATATTTACATATA 10015

RESULT 15  
US-08-187-453-35/c  
Sequence 35, Application US/08187453  
Patent No. 5753431  
GENERAL INFORMATION:  
APPLICANT: Chiang, John  
TITLE OF INVENTION: Cholesterol 7a-Hydroxylase Gene  
TITLE OF INVENTION: Regulatory Elements and Transcription Factors  
NUMBER OF SEQUENCES: 37  
CORRESPONDENCE ADDRESS:  
ADDRESSEE: Foley & Lardner  
STREET: 3000 K Street, N.W., Suite 500  
CITY: Washington, D.C.  
COUNTRY: USA  
ZIP: 20007-5109  
COMPUTER READABLE FORM:  
MEDIUM TYPE: Floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS

